

k__Unclassified_OTU_211; Salmonella -	5.3	6.1	6.6
k__Unclassified_OTU_218; k__Unclassified_OTU_218 -	4.5	5.2	5.7
k__Unclassified_OTU_157; k__Unclassified_OTU_157 -	3.3	2.6	2.2
k__Unclassified_OTU_175; k__Unclassified_OTU_175 -	2.4	2.2	1.7
k__Unclassified_OTU_217; k__Unclassified_OTU_217 -	2	1.7	1.7
k__Unclassified_OTU_136; k__Unclassified_OTU_136 -	1.6	1.7	1.7
k__Unclassified_OTU_87; Bacillus -	2	1.3	1.7
k__Unclassified_OTU_182; k__Unclassified_OTU_182 -	1.6	1.3	1.7
k__Unclassified_OTU_111; Listeria -	0.8	1.3	2.2
k__Unclassified_OTU_40; Lactobacillus -	1.2	1.7	1.3
k__Unclassified_OTU_208; k__Unclassified_OTU_208 -	0.8	1.7	1.3
k__Unclassified_OTU_185; k__Unclassified_OTU_185 -	1.2	1.3	1.3
k__Unclassified_OTU_220; k__Unclassified_OTU_220 -	1.2	1.3	1.3
k__Unclassified_OTU_60; k__Unclassified_OTU_60 -	1.2	1.3	1.3
k__Unclassified_OTU_133; k__Unclassified_OTU_133 -	1.6	0.9	1.3
k__Unclassified_OTU_179; k__Unclassified_OTU_179 -	0.8	0.9	1.3
k__Unclassified_OTU_89; k__Unclassified_OTU_89 -	0.8	0.9	1.3
k__Unclassified_OTU_140; Staphylococcus -	0.8	1.3	0.9
k__Unclassified_OTU_231; k__Unclassified_OTU_231 -	0.8	1.3	0.9
k__Unclassified_OTU_36; k__Unclassified_OTU_36 -	0.8	1.3	0.9
k__Unclassified_OTU_82; k__Unclassified_OTU_82 -	1.2	0.9	0.9
k__Unclassified_OTU_165; k__Unclassified_OTU_165 -	0.8	0.4	1.3
k__Unclassified_OTU_106; Staphylococcus -	0.8	0.9	0.9
k__Unclassified_OTU_162; k__Unclassified_OTU_162 -	0.8	0.9	0.9
k__Unclassified_OTU_173; k__Unclassified_OTU_173 -	0.8	0.9	0.9
	sample1 -	sample2 -	sample3 -

Rarefaction curve

